SEQUENCE LISTING

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<110> Michael F. Thomashow
      Sarah George Fowler
      Jonathan Vogel
      Daniel Zarka
<120> Transcription Factors to Improve Plant Stress Tolerance
<130> 21835-00004
<150> 60/400,777
<151> August 2, 2002
<160> 2
<210> 1
<211> 1308
<212> DNA
<213> Arabidopsis thaliana
<220>
<221>
<222>
<223> This is a cDNA sequence derived from mRNA encoding the putative AP2 domain
transcription factor RAV1 (locus tag: Atlg13260). The RAV1 coding sequence
consists of nucleotides 92 through 1126 as numbered below.
<400> 1
gtatacatat acacaacata attcacaaca caacacaaac acatttctqt tttctccatt
                                                                    60
gtttcaaacc ataaaaaaaa acacagatta aatggaatcg agtagcgttg atgagagtac
                                                                    120
tacaagtaca ggttccatct gtgaaacccc ggcgataact ccggcgaaaa agtcgtcggt
                                                                    180
aggtaactta tacaggatgg gaagcggatc aagcgttgtg ttagattcag agaacggcgt
                                                                    240
agaagctgaa tctaggaagc ttccgtcgtc aaaatacaaa ggtgtggtgc cacaaccaaa
                                                                    300
cggaagatgg ggagctcaga tttacgagaa acaccagcgc gtgtggctcg ggacattcaa
                                                                    360
cgaagaagac gaagccgctc gtgcctacga cgtcgcggtt cacaggttcc gtcqccgtga
                                                                    420
cgccgtcaca aatttcaaag acgtgaagat ggacgaagac gaggtcgatt tcttgaattc
                                                                    480
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tcattcgaaa tctgagatcg ttgatatgtt gaggaaacat acttataacg aagagttaga
                                                                       540
gcagagtaaa cggcgtcgta atggtaacgg aaacatgact aggacgttgt taacgtcggg
                                                                       600
gttgagtaat gatggtgttt ctacgacggg gtttagatcg gcggaggcac tgtttgagaa
                                                                       660
agcggtaacg ccaagcgacq ttgggaaqct aaaccgtttg gttataccga aacatcacqc
                                                                       720
agagaaacat tttccqttac cqtcaaqtaa cqtttccqtq aaaqqaqtqt tqttqaactt
                                                                       780
tgaggacgtt aacgggaaag tgtggaggtt ccgttactcg tattggaaca gtagtcagag
                                                                       840
ttatgttttg actaaaggtt ggagcaggtt cgttaaggag aagaatctac gtgctggtga
                                                                       900
cgtggttagt ttcagtagat ctaacggtca ggatcaacag ttgtacattg ggtggaagtc
                                                                       960
gagatccggg tcagatttag atgcgggtcg ggttttgaga ttgttcggag ttaacatttc
                                                                       1020
accggagagt tcaagaaacg acgtcgtagg aaacaaaaga gtgaacgata ctgagatgtt
                                                                       1080
atcqttqqtq tqtaqcaaqa aqcaacqcat ctttcacqcc tcqtaacaac tcttcttctt
                                                                       1140
ttttttttct tttgttgttt taataatttt taaaaactcc attttcgttt tctttatttg
                                                                       1200
categgtttc tttcttcttg tttaccaaag gttcatgagt tgtttttgtt gtattgatga
                                                                       1260
actgtaaatt ttatttatag gataaatttt aaaaagggtt acttagat
                                                                       1308
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<210> 2
<211> 1308
<212> DNA
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<213> Arabidopsis thaliana

<220>

<221>

<222>

<223> This is a cDNA sequence derived from mRNA encoding the putative zinc-finger protein Zat12 (locus tag: At5g59820). The Zat12 coding sequence consists of nucleotides 102 through 590 as numbered below.

<400> 2

atcatcacaa	ctactatcac	accaaactca	aaaaacacaa	accacaagag	gatcatttca	60
ttttttattg	tttcgtttta	atcatcatca	tcagaagaaa	aatggttgcg	atatcggaga	120
tcaagtcgac	ggtggatgtc	acggcggcga	attgtttgat	gcttttatct	agagttggac	180
		gatcaaaaac				240
agtttcattc	gttccaagcc	ttaggaggtc	accgtgcgag	tcacaagaag	cctaacaacg	300
acgctttgtc	gtctggattg	atgaagaagg	tgaaaacgtc	gtcgcatcct	tgtcccatat	360
gtggagtgga	gtttccgatg	ggacaagctt	tgggaggaca	catgaggaga	cacaggaacg	420
agagtggggc	tgctggtggc	gcgttggtta	cacgcgcttt	gttgccggag	cccacggtga	480
ctacgttgaa	gaaatctagc	agtgggaaga	gagtggcttg	tttggatctg	agtctaggga	540
tggtggacaa	tttgaatctc	aagttggagc	ttggaagaac	agtttattga	ttttatttat	600
tttccttaaa	ttttctgaat	atatttgttt	ctctcattct	ttgaattttt	cttaatattc	660
tagattatac	atacatccgc	agatttagga	aactttcata	gagtgtaatc	ttttctttct	720
gtaaaaatat	attttacttg	tagcattgga	gatttgttat	gagattatct	tacttagcat	780
ttagtgaata	atctattagc	ctattttgcc	gacgtg			816